

SEQUENCE LISTING

<110> Conklin, Darrell C.
Haldeman, Betty A.

<120> MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10

<130> 97-72

<150> 09/199,586

<151> 1998-11-25

<150> 60/066,597

<151> 1997-11-26

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<212> DNA

<213> Homo sapiens

<220>

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<222> (45)...(572)

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Ser Leu Ala Phe Ser Leu Leu Ser Ala Ala Phe Tyr Leu Leu Trp Thr
5 10 15 20

cct tcc act gga ctg aag aca ctc aat ttg gga agc tgt gtg atc gcc 152
Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala
25 30 35

aca aac ctt cag gaa ata cga aat gga ttt tct gac ata cgg ggc agt 200

Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser	
40 45 50	
gtg caa gcc aaa gat gga aac att gac atc aga atc tta agg agg act	248
Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr	
55 60 65	
gag tct ttg caa gac aca aag cct gcg aat cga tgc tgc ctc ctg cgc	296
Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg	
70 75 80	
cat ttg cta aga ctc tat ctg gac agg gta ttt aaa aac tac cag acc	344
His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr	
85 90 95 100	
cct gac cat tat act ctc cgg aag atc agc agc ctc gcc aat tcc ttt	392
Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe	
105 110 115	
ctt acc atc aag aag gac ctc cgg ctc tgt cat gcc cac atg aca tgc	440
Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys	
120 125 130	
cat tgt ggg gag gaa gca atg aag aaa tac agc cag att ctg agt cac	488
His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His	
135 140 145	
ttt gaa aag ctg gaa cct cag gca gca gtt gtg aag gct ttg ggg gaa	536
Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu	
150 155 160	
cta gac att ctt ctg caa tgg atg gag gag aca gaa taggaggaaa	582
Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu	
165 170 175	
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aggcatgacc ccaaaccacc atctctttac tgtactagtc ttgtgctggt cacagtgtat	702
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gagaccatac ttgtataaga tttttgtaat atctttctgc tattggatat atttattagt	822
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<211> 176
 <212> PRT
 <213> Homo sapiens

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			20					25					30		
Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp
		35					40					45			
Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile
	50					55					60				
Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys
65					70					75				80	
Cys	Leu	Leu	Arg	His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys
				85					90					95	
Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
			100						105				110		
Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	His	Ala
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	130					135					140				
Ile	Leu	Ser	His	Phe	Glu	Lys	Leu	Glu	Pro	Gln	Ala	Ala	Val	Val	Lys
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Ser 5	Leu	Ala	Phe	Ser	Leu	Leu	Ser	Ala	Ala	Phe	Tyr	Leu	Leu	Trp	Thr	20					
cct	tcc	act	gga	ctg	aag	aca	ctc	aat	ttg	gga	agc	tgt	gtg	atc	gcc	152					
Pro	Ser	Thr	Gly	Leu	Lys	Thr	Leu	Asn	Leu	Gly	Ser	Cys	Val	Ile	Ala						
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Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp	Ile	Arg	Gly	Ser						
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gtg	caa	gcc	aaa	gat	gga	aac	att	gac	atc	aga	atc	tta	agg	agg	act	248					
Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile	Leu	Arg	Arg	Thr						
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gag	tct	ttg	caa	gac	aca	aag	cct	gcg	aat	cga	tgc	tgc	ctc	ctg	cgc	296					
Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys	Cys	Leu	Leu	Arg						
				70			75				80										
cat	ttg	cta	aga	ctc	tat	ctg	gac	agg	gta	ttt	aaa	aac	tac	cag	acc	344					
His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys	Asn	Tyr	Gln	Thr						
				85			90				95				100						
cct	gac	cat	tat	act	ctc	cgg	aag	atc	agc	agc	ctc	gcc	aat	tcc	ttt	392					
Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu	Ala	Asn	Ser	Phe						
				105			110				115										
ctt	acc	atc	aag	aag	gac	ctc	cgg	ctc	tgt	ctg	gaa	cct	cag	gca	gca	440					
Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	Leu	Glu	Pro	Gln	Ala	Ala						
				120			125				130										
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Val	Val	Lys	Ala	Leu	Gly	Glu	Leu	Asp	Ile	Leu	Leu	Gln	Trp	Met	Glu						
				135			140				145										
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Glu	Thr	Glu																			
				150																	
aagagctcca	gtcttcaata				cctgcagagg				aggcatgacc				ccaaaccacc				atctcttttac	597			
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<211> 151

<212> PRT

<213> Homo sapiens

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Leu	Leu	Trp	Thr	Pro	Ser	Thr	Gly	Leu	Lys	Thr	Leu	Asn	Leu	Gly	Ser
		20					25					30			
Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp
		35				40						45			
Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile
	50				55					60					
Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys
65				70				75						80	
Cys	Leu	Leu	Arg	His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys
			85					90					95		
Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
		100					105					110			
Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	Leu	Glu
	115					120				125					
Pro	Gln	Ala	Ala	Val	Val	Lys	Ala	Leu	Gly	Glu	Leu	Asp	Ile	Leu	Leu
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<213> Homo sapiens

<400> 6

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<211> 45

<212> DNA

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<211> 747

<212> DNA

<213> Homo sapiens

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tgtcatgccc acatgacatg ccattgtggg gaggaagcaa tgaagaaata cagccagatt	300
ctgagtcact ttgaaaagct ggaacctcag gcagcagttg tgaaggcttt gggggaacta	360
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aatattcgag gtcaagagct ccagtccttca atacctgcag aggaggcatg accccaaacc 480
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tgcttccttg catgattgtc tttatgcata cccaatctta attgagacca tacttgtata 600
agatTTTTgt aatatctttc tgctattgga tataatttatt agttaatata tttatttatt 660
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gagctccagt cttcaatacc tgcagaggag gcatgacccc aaaccacat ctctttactg 420
tactagtctt gtgctggtca cagtgtatct tatttatgca ttacttgctt ccttgcata 480
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Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr Glu Ser Leu Gln
35     40     45
Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg His Leu Leu Arg
50     55     60
Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr
65     70     75     80
Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys
85     90     95

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Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu
 100 105 110
 Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu
 115 120 125
 Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu
 130 135 140
 Leu Gln Trp Met Glu Glu Thr Glu
 145 150

<210> 13
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 <213> Homo sapiens

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 Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr Glu Ser Leu Gln
 35 40 45
 Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg His Leu Leu Arg
 50 55 60
 Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr
 65 70 75 80
 Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys
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 100 105 110
 Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu
 115 120 125

<210> 14
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 <213> Homo sapiens

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<210> 15
 <211> 15

<213> Homo sapiens

<400> 15

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<211> 15

<212> PRT

<213> Homo sapiens

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<211> 15

<212> PRT

<213> Homo sapiens

<400> 17

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<210> 18

<211> 824

<212> DNA

<213> Mus musculus

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<222> (71)...(598)

<400> 18

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      Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala
          1             5             10

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Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His
15 20 25

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ttt tct gag att cgg gat agt gtg caa gct gaa gat aca aat att gac Phe Ser Glu Ile Arg Asp Ser Val Gln Ala Glu Asp Thr Asn Ile Asp 50 55 60	253
atc aga att tta agg acg act gag tct ttg aaa gac ata aag tct ttg Ile Arg Ile Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu 65 70 75	301
gat agg tgc tgc ttc ctt cgt cat cta gtg aga ttc tat ctg gac agg Asp Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg 80 85 90	349
gta ttc aaa gtc tac cag acc cct gac cac cat acc ctg aga aag atc Val Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile 95 100 105	397
agc agc ctc gcc aac tcc ttt ctt atc atc aag aag gac ctc tca gtc Ser Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val 110 115 120 125	445
tgt cat tct cac atg gca tgt cat tgt ggg gaa gaa gca atg gag aaa Cys His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys 130 135 140	493
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gag atg cta tagatgaaag tggagaggct gctgagaaca ctcctgtcca Glu Met Leu 175	638
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<211> 176

<212> PRT

<213> Mus musculus

<400> 19

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Leu	Leu	Trp	Thr	Pro	Leu	Thr	Gly	Leu	Lys	Thr	Leu	His	Leu	Gly	Ser
			20				25					30			
Cys	Val	Ile	Thr	Ala	Asn	Leu	Gln	Ala	Ile	Gln	Lys	Glu	Phe	Ser	Glu
	35					40				45					
Ile	Arg	Asp	Ser	Val	Gln	Ala	Glu	Asp	Thr	Asn	Ile	Asp	Ile	Arg	Ile
	50				55				60						
Leu	Arg	Thr	Thr	Glu	Ser	Leu	Lys	Asp	Ile	Lys	Ser	Leu	Asp	Arg	Cys
65				70				75				80			
Cys	Phe	Leu	Arg	His	Leu	Val	Arg	Phe	Tyr	Leu	Asp	Arg	Val	Phe	Lys
			85					90				95			
Val	Tyr	Gln	Thr	Pro	Asp	His	His	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
			100					105				110			
Ala	Asn	Ser	Phe	Leu	Ile	Ile	Lys	Lys	Asp	Leu	Ser	Val	Cys	His	Ser
	115						120					125			
His	Met	Ala	Cys	His	Cys	Gly	Glu	Glu	Ala	Met	Glu	Lys	Tyr	Asn	Gln
	130					135				140					
Ile	Leu	Ser	His	Phe	Ile	Glu	Leu	Glu	Leu	Gln	Ala	Ala	Val	Val	Lys
145				150				155				160			
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<212> PRT

<213> Mus musculus

<400> 20

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<213> Mus musculus

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1 5 10 15

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 35 40 45
 Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys
 50 55 60
 Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu
 65 70 75 80
 Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser
 85 90 95
 His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln
 100 105 110
 Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys
 115 120 125
 Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu
 130 135 140

<210> 26
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 <213> Homo sapiens

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 1 5 10 15

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 Leu Arg Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys
 35 40 45
 Cys Leu Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys
 50 55 60
 Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu
 65 70 75 80
 Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala
 85 90 95
 His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln
 100 105 110
 Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys
 115 120 125
 Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu Glu Thr Glu
 130 135 140

<210> 27

<211> 38

<212> PRT

<213> Homo sapiens

<400> 27

Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe
 1 5 10 15

Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu
 20 25 30
 Asp Ile Leu Leu Gln Trp
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<210> 28

<211> 71

<212> PRT

<213> Homo sapiens

<400> 28

Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg
 1 5 10 15
 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg
 20 25 30
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu
 35 40 45

Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr
 50 55 60
 Gln Thr Pro Asp His Tyr Thr
 65 70

<210> 29
 <211> 92
 <212> PRT
 <213> Homo sapiens

<400> 29
 Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg
 1 5 10 15
 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg
 20 25 30
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu
 35 40 45
 Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr
 50 55 60
 Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn
 65 70 75 80
 Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys
 85 90

<210> 30
 <211> 82
 <212> PRT
 <213> Homo sapiens

<400> 30
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu
 1 5 10 15
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp
 20 25 30
 Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu Glu Ala
 35 40 45
 Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro
 50 55 60
 Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln
 65 70 75 80
 Trp Met

<210> 31

<211> 36
 <212> PRT
 <213> Homo sapiens

<400> 31
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu
 1 5 10 15
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp
 20 25 30
 Leu Arg Leu Cys
 35

<210> 32
 <211> 61
 <212> PRT
 <213> Homo sapiens

<400> 32
 Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His
 1 5 10 15
 Ala His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser
 20 25 30
 Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val
 35 40 45
 Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met
 50 55 60

<210> 33
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<220>
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 <222> (71)...(532)

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 Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala
 1 5 10
 gtg ggt ttt ctt ctc tgg act cct tta act ggg ctc aag acc ctc cat 157

Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His	
15 20 25	
ttg gga agc tgt gtg att act gca aac cta cag gca ata caa aag gaa	205
Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu	
30 35 40 45	
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Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu	
50 55 60	
cgt cat cta gtg aga ttc tat ctg gac agg gta ttc aaa gtc tac cag	301
Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln	
65 70 75	
acc cct gac cac cat acc ctg aga aag atc agc agc ctc gcc aac tcc	349
Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser	
80 85 90	
ttt ctt atc atc aag aag gac ctc tca gtc tgt cat tct cac atg gca	397
Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala	
95 100 105	
tgt cat tgt ggg gaa gaa gca atg gag aaa tac aac caa att ctg agt	445
Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser	
110 115 120 125	
cac ttc ata gag ttg gaa ctt cag gca gcg gtg gta aag gct ttg gga	493
His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly	
130 135 140	
gaa cta ggc att ctt ctg aga tgg atg gag gag atg cta tagatgaaag	542
Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu	
145 150	
tggaataggct gctgagaaca ctctgtcca agaattctcag acctcagcac catgaagaca	602
tggtcccccagg tgctggcatt tctactcaag agttccagtc ctcagcacca cgaagatggc	662
ctcaaaccac caccctttg tgatataact tagtgctagc tatgtgtata ttatttctac	722
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<210> 34

<211> 154

<212> PRT

<213> Mus musculus

<400> 34

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Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala Val Gly Phe
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Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His Leu Gly Ser
 20           25           30
Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu Phe Ser Glu
 35           40           45
Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu Arg His Leu
 50           55           60
Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln Thr Pro Asp
 65           70           75           80
His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Ile
 85           90           95
Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala Cys His Cys
 100          105          110
Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser His Phe Ile
 115          120          125
Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Gly
 130          135          140
Ile Leu Leu Arg Trp Met Glu Glu Met Leu
145          150

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<210> 35

<211> 130

<212> PRT

<213> Mus musculus

<400> 35

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Leu Lys Thr Leu His Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln
 1           5           10           15

Ala Ile Gln Lys Glu Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp
 20           25           30
Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val
 35           40           45
Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser
 50           55           60
Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys
 65           70           75           80
His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr

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				85					90					95	
Asn	Gln	Ile	Leu	Ser	His	Phe	Ile	Glu	Leu	Glu	Leu	Gln	Ala	Ala	Val
			100					105					110		
Val	Lys	Ala	Leu	Gly	Glu	Leu	Gly	Ile	Leu	Leu	Arg	Trp	Met	Glu	Glu
		115					120					125			
Met	Leu														
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<210> 36
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<400> 36
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27

<210> 37
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<400> 37
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17

<210> 38
 <211> 25
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 <213> Mus musculus

<400> 38
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25

<210> 39
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<400> 39
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26

<210> 41

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<400> 41

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ggtgcatatt cctgggtggct aga

23

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<212> DNA

<213> Mus musculus

<400> 43

attgcagtgt aagggaatac agaga

25